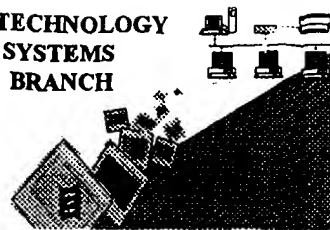


RAW SEQUENCE LISTING ERROR REPORT

BIOTECHNOLOGY
SYSTEMS
BRANCH



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

09/489,760B

Source:

1600

Date Processed by STIC:

10-29-03

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to (EFFECTIVE 12/01/2003):
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/2003

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER:

09/489,760 B

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use **space characters**, instead.
- 4 Non-ASCII The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. Please **ensure your subsequent submission is saved in ASCII text.**
- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**
- 7 Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for **each** skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.
- 8 Skipped Sequences
 (NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000
- 9 Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is **MANDATORY** if n's or Xaa's are present.
 In <220> to <223> section, please explain location of **n** or **Xaa**, and which residue **n** or **Xaa** represents.
- 10 X Invalid <213>
 Response Per 1.823 of Sequence Rules, the only **valid** <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence
- 11 Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.
 Use of <220> to <223> is **MANDATORY** if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 00/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n/Xaa "**n**" can only represent a single nucleotide; "**Xaa**" can only represent a single amino acid



1600

RAW SEQUENCE LISTING

DATE: 10/29/2003

PATENT APPLICATION: US/09/489,760B

TIME: 10:35:51

Input Set : A:\SEQLIST.CRF.txt

Output Set: N:\CRF4\10292003\I489760B.raw

3 <110> APPLICANT: Goulmy, Elsa A.J.M
 4 Hunt, Donald F
 5 Engelhard, Victor H
 7 <120> TITLE OF INVENTION: The HA-1 Antigen
 9 <130> FILE REFERENCE: 2183-4285US
 11 <140> CURRENT APPLICATION NUMBER: 09/489,760B
 12 <141> CURRENT FILING DATE: 2000-01-21
 14 <150> PRIOR APPLICATION NUMBER: PCT/NL98/00424
 15 <151> PRIOR FILING DATE: 1998-07-23
 17 <160> NUMBER OF SEQ ID NOS: 17
 19 <170> SOFTWARE: PatentIn Ver. 2.1
 21 <210> SEQ ID NO: 1
 22 <211> LENGTH: 9
 23 <212> TYPE: PRT
 24 <213> ORGANISM: Histocompatibility antigen
 26 <220> FEATURE:
 27 <221> NAME/KEY: MISC_FEATURE
 28 <222> LOCATION: (3)..(3)
 29 <223> OTHER INFORMATION: Xaa is HISTIDINE OR ARGININE RESIDUE
 31 <400> SEQUENCE: 1

**Does Not Comply
Corrected Diskette Needed**

*see item 10 on error
summary report.*

W--> 32 Val Leu Xaa Asp Asp Leu Leu Glu Ala

33 1 5
 36 <210> SEQ ID NO: 2
 37 <211> LENGTH: 9
 38 <212> TYPE: PRT
 39 <213> ORGANISM: Histocompatibility antigen

41 <400> SEQUENCE: 2
 42 Val Leu His Asp Asp Leu Leu Glu Ala
 43 1 5

46 <210> SEQ ID NO: 3
 47 <211> LENGTH: 9
 48 <212> TYPE: PRT
 49 <213> ORGANISM: Histocompatibility antigen

51 <220> FEATURE:
 52 <221> NAME/KEY: MISC_FEATURE
 53 <222> LOCATION: (2)..(2)
 54 <223> OTHER INFORMATION: Xaa is ISOLEUCINE OR LEUCINE
 56 <400> SEQUENCE: 3

W--> 57 Tyr Xaa Thr Asp Arg Val Met Thr Val

58 1 5
 61 <210> SEQ ID NO: 4
 62 <211> LENGTH: 9
 63 <212> TYPE: PRT

RAW SEQUENCE LISTING

DATE: 10/29/2003

PATENT APPLICATION: US/09/489,760B

TIME: 10:35:51

Input Set : A:\SEQLIST.CRF.txt

Output Set: N:\CRF4\10292003\I489760B.raw

64 <213> ORGANISM: Histocompatibility antigen

66 <220> FEATURE:

67 <221> NAME/KEY: MISC_FEATURE

68 <222> LOCATION: (2)..(7)

69 <223> OTHER INFORMATION: Xaa is LEUCINE OR ISOLEUCINE

71 <400> SEQUENCE: 4

W--> 72 Val Xaa His Asp Asp Xaa Xaa Glu Ala

73 1 5

76 <210> SEQ ID NO: 5

77 <211> LENGTH: 9

78 <212> TYPE: PRT

79 <213> ORGANISM: KIAA0723 partial complementary DNA

81 <400> SEQUENCE: 5

82 Val Leu Arg Asp Asp Leu Leu Glu Ala

83 1 5

86 <210> SEQ ID NO: 6

87 <211> LENGTH: 25

88 <212> TYPE: DNA

89 <213> ORGANISM: Unknown

91 <220> FEATURE:

92 <223> OTHER INFORMATION: Reverse primer

94 <400> SEQUENCE: 6

95 gctctctgcat gacgcctctgt ctgca 25

98 <210> SEQ ID NO: 7

99 <211> LENGTH: 24

100 <212> TYPE: DNA

101 <213> ORGANISM: Unknown

103 <220> FEATURE:

104 <223> OTHER INFORMATION: Forward primer

106 <400> SEQUENCE: 7

107 gacgtcgtcg aggacatctc ccat 24

110 <210> SEQ ID NO: 8

111 <211> LENGTH: 25

112 <212> TYPE: DNA

113 <213> ORGANISM: Unknown

115 <220> FEATURE:

116 <223> OTHER INFORMATION: Reverse Primer

118 <400> SEQUENCE: 8

119 gaaggccaca gcaatcgtct ccagg 25

122 <210> SEQ ID NO: 9

123 <211> LENGTH: 30

124 <212> TYPE: DNA

125 <213> ORGANISM: Unknown

127 <220> FEATURE:

128 <223> OTHER INFORMATION: HA-1" allele primer H1

130 <400> SEQUENCE: 9

131 ccttgagaaa cttaaggagt gtgtgctgca 30

134 <210> SEQ ID NO: 10

135 <211> LENGTH: 30

same error

RAW SEQUENCE LISTING

DATE: 10/29/2003

PATENT APPLICATION: US/09/489,760B

TIME: 10:35:51

Input Set : A:\SEQLIST.CRF.txt

Output Set: N:\CRF4\10292003\I489760B.raw

```

136 <212> TYPE: DNA
137 <213> ORGANISM: Unknown
139 <220> FEATURE:
140 <223> OTHER INFORMATION: HA-1r allele primer R1
142 <400> SEQUENCE: 10
143 ccttgagaaa cttaaggagt gtgtgttgcg          30
146 <210> SEQ ID NO: 11
147 <211> LENGTH: 33
148 <212> TYPE: DNA
149 <213> ORGANISM: Forward KIAA00223 based PCR primer
151 <400> SEQUENCE: 11
152 ccggcatgga cgtcctcgag gacatctccc atc          33
155 <210> SEQ ID NO: 12
156 <211> LENGTH: 30
157 <212> TYPE: DNA
158 <213> ORGANISM: Reverse KIAA00223 based PCR primer
160 <400> SEQUENCE: 12
161 ctacttcagg ccacagcaat cgtctccagg          30
164 <210> SEQ ID NO: 13
165 <211> LENGTH: 38
166 <212> TYPE: DNA
167 <213> ORGANISM: Unknown
169 <220> FEATURE:
170 <223> OTHER INFORMATION: HA-1-/-
172 <400> SEQUENCE: 13
173 gagtgtgtgt tgcgtgacga cctccttgag gcccgcgg          38
176 <210> SEQ ID NO: 14
177 <211> LENGTH: 13
178 <212> TYPE: PRT
179 <213> ORGANISM: Unknown
181 <220> FEATURE:
182 <223> OTHER INFORMATION: HA-1-/-
184 <400> SEQUENCE: 14
185 Glu Cys Val Leu Arg Asp Asp Leu Leu Glu Ala Arg Arg
186 1          5          10
189 <210> SEQ ID NO: 15
190 <211> LENGTH: 38
191 <212> TYPE: DNA
192 <213> ORGANISM: Unknown
194 <220> FEATURE:
195 <223> OTHER INFORMATION: HA-1+/+
197 <400> SEQUENCE: 15
198 gagtgtgtgc tgcgtgacga cctccttgag gcccgcgg          38
201 <210> SEQ ID NO: 16
202 <211> LENGTH: 13
203 <212> TYPE: PRT
204 <213> ORGANISM: Unknown
206 <220> FEATURE:
207 <223> OTHER INFORMATION: HA-1+/+

```

RAW SEQUENCE LISTING

PATENT APPLICATION: **US/09/489,760B**

DATE: 10/29/2003

TIME: 10:35:51

Input Set : **A:\SEQLIST.CRF.txt**

Output Set: **N:\CRF4\10292003\I489760B.raw**

```

209 <400> SEQUENCE: 16
210 Glu Cys Val Leu His Asp Asp Leu Leu Glu Ala Arg Arg
211 1          5          10
214 <210> SEQ ID NO: 17
215 <211> LENGTH: 10
216 <212> TYPE: PRT
217 <213> ORGANISM: Unknown
219 <220> FEATURE:
220 <223> OTHER INFORMATION: Peptide that can bind recombinant HLA-A2.1 and
b2microglobulin
222 <400> SEQUENCE: 17
223 Phe Leu Pro Ser Asp Cys Phe Pro Ser Val
224 1          5          10

```

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/489,760B

DATE: 10/29/2003
TIME: 10:35:52

Input Set : A:\SEQLIST.CRF.txt
Output Set: N:\CRF4\10292003\I489760B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 3
Seq#:3; Xaa Pos. 2
Seq#:4; Xaa Pos. 2,6,7

VERIFICATION SUMMARY

DATE: 10/29/2003

PATENT APPLICATION: US/09/489,760B

TIME: 10:35:52

Input Set : A:\SEQLIST.CRF.txt

Output Set: N:\CRF4\10292003\I489760B.raw

L:32 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0

L:57 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0

L:72 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0